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REMARKS

In compliance with 37 C.F.R. § 1.821 through § 1.825, applicants have amended the specification to add Sequence Identifiers. No new matter has been added by this amendment. Applicants respectfully request entry of the present amendment.

Attached hereto is a marked up version of the changes made to the specification by the current amendment with additions underlined and deletions bracketed. The attached pages are captioned "**VERSION WITH MARKINGS TO SHOW CHANGES MADE**".

CONCLUSION

In the unlikely event that the transmittal letter is separated from this document and the Patent Office determines that an extension and/or other relief is required, Applicants petition for any required relief including extensions of time and authorize the Assistant Commissioner to charge the cost of such petitions and/or other fees due in connection with the filing of this document to **Deposit Account No. 03-1952** referencing docket no. 348022001600. However, the Assistant Commissioner is not authorized to charge the cost of the issue fee to the Deposit Account.

Respectfully submitted,

Dated: October 26, 2001

By: _____

Debra J. Glaister
Registration No. 33,888

Morrison & Foerster LLP
755 Page Mill Road
Palo Alto, California 94304-1018
Telephone: (650) 813-5725
Facsimile: (650) 494-0792

VERSION WITH MARKINGS TO SHOW CHANGES MADE

METHODS OF TREATING NEOPLASIA WITH COMBINATION TARGET CELL-SPECIFIC ADENOVIRUS, CHEMOTHERAPY AND RADIATION

On page 17, the paragraph containing line 20 has been amended, as follows:;

Figure 37 depicts a nucleotide SEQ ID NO: 17 and amino acid sequence SEQ ID NO: 18 for ADP.

On page 77, please amend the paragraph containing lines 28 and 31, as follows:

In some embodiments, a melanocyte-specific TRE comprises sequences derived from the 5' flanking region of a human tyrosinase gene depicted in Table 14. In some of these embodiments, the melanocyte-specific TRE comprises tyrosinase nucleotides from about -231 to about +65 relative to the transcription start site (from about nucleotide 244 to about nucleotide 546 of SEQ ID NO:[__]10) and may further comprise nucleotides from about -1956 to about -1716 relative to the human tyrosinase transcription start site (from about nucleotide 6 to about nucleotide -243 of SEQ ID NO:[__]10). A tyrosinase TRE can comprise nucleotides from about -231 to about + 65 juxtaposed to nucleotides from about -1956 to about -1716. It has been reported that nucleotides from about -1956 to about -1716 relative to the human tyrosinase transcription start site can confer melanocyte-specific expression of an operably linked reporter gene with either a homologous or a heterologous promoter. Accordingly, in some embodiments, a melanocyte-specific TRE comprises nucleotides from about -1956 to about -1716 operably linked to a heterologous promoter.

On page 86, the paragraph containing lines 30 and 33 has been amended, as follows:

IRES elements were first discovered in picornavirus mRNAs (Jackson RJ, Howell MT, Kaminski A (1990) *Trends Biochem Sci* 15(12):477-83) and Jackson RJ and Kaminski, A. (1995) *RNA* 1(10):985-1000). The present invention provides improved adenovirus vectors comprising co-transcribed first and second genes under transcriptional control of a heterologous, target cell-specific TRE, and wherein the second gene (i.e., coding region) is under translational control of an internal ribosome entry site (IRES). Any IRES may be used in the adenovirus vectors of the invention, as long as they exhibit requisite function in the vectors. Example of IRES which can be used in the present invention include those provided in Table I and referenced in Table II.

Examples of IRES elements include the encephelomycarditis virus (EMCV) which is commercially available from Novagen (Duke et al. (1992) *J. Virol* 66(3):1602-9) the sequence for which is depicted in Table 12 (SEQ ID NO:1). Another example of an IRES element disclosed herein is the VEGF IRES (Huez et al. (1998) *Mol Cell Biol* 18(11):6178-90). This IRES has a short segment and the sequence is depicted in Table 12 (SEQ ID NO:2).

On page 131, Table 9 has been amended, as follows:

Table 9

Primer	Sequence	Note
A.	5'-GACGTCGACTAATTCCGGTTATTTTCCA SEQ ID NO: 19	For PCR EMCV IRES, *GTCGAC* is a SalI site.
B.	5'-GACGTCGACATCGTGTTTTTCAAAGGAA SEQ ID NO: 20	For PCR EMCV IRES, *GTCGAC* is a SalI site.
C.	5'-CCTGAGACGCCCGACATCACCTGTG SEQ ID NO: 21	Ad5 sequence to 1314 to 1338.
D.	5'-GTCGACCATTCAGCAAACAAAGGCGTTAAC SEQ ID NO: 22	Antisense of Ad5 sequence 1572 to 1586. *GTCGAC* is a SalI site. Underline region overlaps with E.
E.	5'-TGCTGAATGGTCGACATGGAGGCTTGGGAG SEQ ID NO: 23	Ad5 sequence 1714 to 1728. *GTCGAC* is a SalI site. Underline region overlaps with D.
F.	5'-CACAAACCGCTCTCCACAGATGCATG SEQ ID NO: 24	Antisense of Ad5 sequence 2070 to 2094.

On page 134, the paragraph containing lines 22 and 23 has been amended, as follows:

The 519 base pair EMCV IRES segment was PCR amplified from Novagen's pCITE vector by primers A/B:

primer A: 5'-GACGTCGACTAATTCCGGTTATTTTCCA SEQ ID NO: 19

primer B 5'-GACGTCGACATCGTGTTTTTCAAAGGAA SEQ ID NO: 20 (*GTCGAC* is a SalI site).

On page 135, the paragraph containing lines 13 and 14 has been amended, as follows:

CP1088

The 2.2kb (-2225 to +1) human UPII was amplified from CP657 with primer 127.2.1 (5'-AGGACCGGTCACTATAGGGCACGCGTGGT-3'(SEQ ID NO: 25)) PLUS 127.2.2 (5'-AGGACCGGTGGGATGCTGGGCTGGGAGGTGG-3'(SEQ ID NO: 26)) and digested with PinAI and ligated with CP629 cut with PinAI.

On page 137, Table 11 has been amended, as follows:

TABLE 11

Name	Vector	Ad 5 Vector	E1A TRE	E1B TRE	E3
CV874	CP1086	pBHGE3	1.9 kb mUPII	IRES	intact
CV875	CP1087	pBHGE3	1.0 kb hUPII	IRES	intact
CV876	CP1088	pBHGE3	2.2 kb hUPII	IRES	intact
CV877	CP1089	pBHGE3	1.0 kb mUPII	1.0 kb hUPII (E1B promoter deleted)	intact
CV882	CP1129	pBHGE3	1.8 kb hUPII	IRES	intact
CV884	CP1131	pBHGE3	1.8 kb hUPii	IRES (E1B 19-kDa deleted)	intact

Viruses are tested and characterized as described above.

Primer sequences:

96.74.1	GACGTCGACATCGTGTTTTTCAAAGGAA SEQ ID NO: 20
96.74.2	GACGTCGACTAATTCCGGTTATTTTCCA SEQ ID NO: 19
96.74.3	CCTGAGACGCCCGACATCACCTGTG SEQ ID NO: 21
96.74.4	TGCTGAATGGTCGACATGGAGGCTTGGGAG SEQ ID NO: 23
96.74.5	CACAACCGCTCTCCACAGATGCATG SEQ ID NO: 24
96.74.6	GTCGACCATTCAGCAAACAAAGGCGTTAAC SEQ ID NO: 22
100.113.1	AGGGGTACCCACTATAGGGCACGCGTGGT SEQ ID NO: 27
100.113.2	ACCCAAGCTTGGGATGCTGGGCTGGGAGGTGG SEQ ID NO: 28
127.2.2	AGGACCGGTGGGATGCTGGGCTGGGAGGTGG SEQ ID NO: 26
127.50.1	AGGACCGGTCAGGCTTCACCCCAGACCCAC SEQ ID NO: 29
31.166.1	TGCGCCGGTGTACACAGGAAGTGA SEQ ID NO: 30

32.32.1	GAGTTTGTGCCATCGGTCTAC SEQ ID NO: 31
32.32.2	AATCAATCCTTAGTCCTCCTG SEQ ID NO: 32
51.176	GCAGAAAAATCTTCCAAACACTCCC SEQ ID NO: 33
99.120.1	ACGTACACCGGTCGTTACATAACTTAC SEQ ID NO: 34
99.120.2	CTAGCAACCGGTCGGTTCACTAAACG SEQ ID NO: 35

On page 139, the paragraph containing line 11 has been amended, as follows:

B. ***Example 16: Construction of a Replication-Competent Adenovirus Vector with a CEA-TRE and a EMCV IRES***

Using a strategy similar to Example 1, the TRE fragment from Carcinembryonic antigen (CEA)(Table 14, SEQ ID NO:[__]14) is used to construct virus designated CV873. A PinAI fragment containing the CEA-TRE was cloned into the PinAI site in front of E1A of CP627 for the transcriptional control. The resultant plasmid CP1080 is used together with pBHGE3 to generate CV873.

On page 167, the paragraph containing line 3 has been amended, as follows:

1. Table 12: IRES Sequences

SEQ ID NO:[__]1 A 519 base pair IRES obtainable from encephelomycarditis virus (EMCV).

```
1     GACGTCGACTAATTCCGGTTATTTTCCACCATATTGCCGTCTTTTGGCAA
          SalI
51    TGTGAGGGCCCGGAAACCTGGCCCTGTCTTCTTGACGAGCATTCCTAGGG
101   GTCTTTCCCCTCTCGCCAAAGGAATGCAAGGTCTGTTGAATGTCGTGAAG
151   GAAGCAGTTCCTCTGGAAGCTTCTTGAAGACAAACAACGTCTGTAGCGAC
201   CCTTTGCAGGCAGCGGAACCCCCCACCTGGCGACAGGTGCCTCTGCGGCC
251   AAAAGCCACGTGTATAAGATACACCTGCAAAGGCGGCACAACCCCAGTGC
301   CACGTTGTGAGTTGGATAGTTGTGGAAAGAGTCAAATGGCTCTCCTCAAG
351   CGTATTCAACAAGGGGCTGAAGGATGCCCAGAAGGTACCCCATTGTATGG
401   GATCTGATCTGGGGCCTCGGTGCACATGCTTTACATGTGTTTAGTCGAGG
```

451 TTAAAAAACGTCTAGGCCCCCCGAACCACGGGGACGTGGTTTTCCTTTGA

SalI

501 AAAACACGAT*GTCGAC*GTC

On page 167, the paragraph containing line 19 has been amended, as follows:

SEQ ID NO:[__]2 An IRES obtainable from vascular endothelial growth factor (VEGF).

1 ACGTAGTCGACAGCGCAGAGGCTTGGGGCAGCCGAGCGGCAGCCAGGCCC

SalI

51 CGGCCCGGGCCTCGGTTCCAGAAGGGAGAGGAGCCCGCCAAGGCGCGCAA

101 GAGAGCGGGCTGCCTCGCAGTCCGAGCCGGAGAGGGAGCGCGAGCCGCGC

151 CGGCCCCGGACGGCCTCCGAAACCATGGTCGACACGTA

SalI

On page 167, the paragraph containing line 28 has been amended, as follows:

SEQ ID NO:[__]3 A 5'UTR region of HCV.

1 GCCAGCCCCCTGATGGGGGCGACACTCCGCCATGAATCACTCCCCTGTGAGGAACTACTG

61 TCTTCACGCAGAAAGCGTCTAGCCATGGCGTTAGTATGAGTGTCGTGCAGCCTCCAGGAC

121 CCCCCCTCCCGGGAGAGCCATAGTGGTCTGCGGAACCGGTGAGTACACCGGAATTGCCAG

181 GACGACCGGGTCCTTTCTTGGATTAACCCGCTCAATGCCTGGAGATTTGGGCGTGCCCCC

241 GCAAGACTGCTAGCCGAGTAGTGTTGGGTCGCGAAAGGCCTTGTGGTACTGCCTGATAGG

301 GTGCTTGCGAGTGCCCCGGGAGGTCTCGTAGACCGTGCACC (341)

On page 168, the paragraph containing 1 has been amended, as follows:

SEQ ID NO:[__]4 A 5'UTR region of BiP SEQ ID NO:4

1 CCCGGGGTCACTCCTGCTGGACCTACTCCGACCCCCTAGGCCGGGAGTGAAGGCGGGACT

61 TGTGCGGTTACCAGCGGAAATGCCTCGGGGTCAGAAGTCGCAGGAGAGATAGACAGCTGC

121 TGAACCAATGGGACCAGCGGATGGGGCGGATGTTATCTACCATTGGTGAACGTTAGAAAC

181 GAATAGCAGCCAATGAATCAGCTGGGGGGGCGGAGCAGTGACGTTTATTGCGGAGGGGGC

241 CGCTTCGAATCGGCGGCGGCCAGCTTGGTGGCCTGGGCCAATGAACGGCCTCCAACGAGC

```
301   AGGGCCTTCACCAATCGGCGGCCTCCACGACGGGGCTGGGGGAGGGTATATAAGCCGAGT

361   AGGCGACGGTGAGGTCGACGCCGGCCAAGACAGCACAGACAGATTGACCTATTGGGGTGT

421   TTCGCGAGTGTGAGAGGGAAGCGCCGCGGCCTGTATTTCTAGACCTGCCCTTCGCCTGGT

481   TCGTGGCGCCTTGTGACCCCGGGCCCCTGCCGCCTGCAAGTCGAAATTGCGCTGTGCTCC

541   TGTGCTACGGCCTGTGGCTGGACTGCCTGCTGCTGCCCAACTGGCTGGCAAGATG (595)
```

On page 168, the paragraph containing line 15 has been amended, as follows:

SEQ ID NO:[__]5 A 5'UTR of PDGF SEQ ID NO:5

```
1     GTTTGCACCTCTCCCTGCCCGGGTGCTCGAGCTGCCGTTGCAAAGCCAACTTTGGAAAAA

61    GTTTTTTGGGGGAGACTTGGGCCTTGAGGTGCCCAGCTCCGCGCTTTCCGATTTTGGGGG

121   CTTTCCAGAAAATGTTGCAAAAAAGCTAAGCCGGCGGGCAGAGGAAAACGCCTGTAGCCG

181   GCGAGTGAAGACGAACCATCGACTGCCGTGTTCCTTTTCCTCTTGGAGGTTGGAGTCCCC

241   TGGGCGCCCCCACACCCCTAGACGCCTCGGCTGGTTCGCGACGCAGCCCCCCGGCCGTGG

301   ATGCTGCACTCGGGCTCGGGATCCGCCCAGGTAGCCGGCCTCGGACCCAGGTCCTGCGCC

361   CAGGTCCTCCCCTGCCCCCCAGCGACGGAGCCGGGGCCGGGGGCGGCGGCGCCGGGGGCA

421   TGCGGGTGAGCCGCGGCTGCAGAGGCCTGAGCGCCTGATCGCCGCGGACCTGAGCCGAGC

481   CCACCCCCCTCCCCAGCCCCCCACCCTGGCCGCGGGGGCGGCGCGCTCGATCTACGCGTC

541   CGGGGCCCCGCGGGGCCGGGCCCGGAGTCGGCATG (575)
```

Beginning on page 169 and ending on page 172, the paragraph containing lines 7 (page 169) and 1 (page 170) has been amended, as follows:

2. Table 14: TRE Sequences

Nucleotide sequence of a human uroplakin II 5' flanking region. Position +1 (the translational start site) is denoted with an asterisk. SEQ ID NO:[__]6 (number 1 of SEQ ID NO:[__]6 corresponds to position -2239 with respect to the translational start site).

```
TCGATAGGTA  CCCACTATAG  GGCACGCGTG  GTCGACGGCC  CGGGCTGGTC
1                                                        50

TGGCAACTTC  AAGTGTGGGC  CTTTCAGACC  GGCATCATCA  GTGTTACGGG
```

51 100

GAAGTCACTA GGAATGCAGA ATTGATTGAG CACGGTGGCT CACACCTGTA
101 150

ATCCCAACAC TCTGGGAGGC CAAGGCAGGT GGATCACTTG TGGTCAGGAG
151 200

TTTGAGACCA GCCTGGCCAA CATGGTGAAA CCTCATCTCT ACTAAAAATA
201 250

CAAAAATTAG CTGGGAATGG TGGCACATGC CTATAATCCC AGTTACTCAG
251 300

GAGGCTGAGG CAGGAGAATC ATTTGAACCT GGGAGGCAGA GGTTGCAGTG
301 350

AGCCGAGATC ACGCCACTGC ACTCCAGCCT GGGTGACACA GCGAGACTCT
351 400

GTCTCAAAAA AAAAAAAATG CAGAATTTCA GGCTTCACCC CAGACCCACT
401 450

GCATGACTGC ATGAGAAGCT GCATCTTAAC AAGATCCCTG GTAATTCATA
451 500

CGCATATTAA ATTTGGAGAT GCACTGGCGT AAGACCCTCC TACTCTCTGC
501 550

TTAGGCCCAT GAGTTCTTCC TTTACTGTCA TTCTCCACTC ACCCCAAACT
551 600

TTGAGCCTAC CCTTCCCACC TTGGCGGTAA GGACACAACC TCCCTCACAT
601 650

TCCTACCAGG ACCCTAAGCT TCCCTGGGAC TGAGGAAGAT AGAATAGTTC
651 700

GTGGAGCAAA CAGATATACA GCAACAGTCT CTGTACAGCT CTCAGGCTTC
701 750

TGGAAGTTCT ACAGCCTCTC CCGACAAAGT ATTCCACTTT CCACAAGTAA
751 800

CTCTATGTGT CTGAGTCTCA GTTTCCACTT TTCTCTCTCT CTCTCTCTCT
801 850

CAACTTTCTG AGACAGAGTT TCACTTAGTC GCCCAGGCTG GAGTGCAGGG
851 900

GCACAATCTC GGCTCACTGC AACCTCCACC TCCTGGGTTC AAGTGTTTCT
901 950

CCTGTCTCAG CCTCCCGAGT AGCTGGGATT ACAGGCACAC ACCACCGCGT

951 1000

TAGTTTTTGT ATTTTTGGTA GAGATGGTGT TTCGCCATAT TGGCCAGGCT
1001 1050

GATCTCGAAC TCCTGACCTC AGGTGATCCG CCCACCTCGG CCTCCCAAAG
1051 1100

TGCTGGGATT ACAGGCATGA GCCACCACGC CCGGCTGATC TCTTTTCTAT
1101 1150

TTTAATAGAG ATCAAACTCT CTGTGTTGCC TAGGCTGGTC TTGAACTCCT
1151 1200

GGCCTCGAGT GATCCTCCCA CCTTGGCCTC CCAAAGTGTT GAGATTACAG
1201 1250

GCATGAGCCA CTGTGCCTGG CCTCAGTTCT ACTACAAAAG GAAGCCAGTA
1251 1300

CCAGCTACCA CCCAGGGTGG CTGTAGGGCT ACAATGGAGC ACACAGAACC
1301 1350

CCTACCCAGG GCCCGGAAGA AGCCCCGACT CCTCTCCCCT CCCTCTGCCC
1351 1400

AGAACTCCTC CGCTTCTTTC TGATGTAGCC CAGGGCCGGA GGAGGCAGTC
1401 1450

AGGGAAGTTC TGTCTCTTTT TCATGTTATC TTACGAGGTC TCTTTTCTCC
1451 1500

ATTCTCAGTC CAACAAATGG TTGCTGCCCA AGGCTGACTG TGCCCACCCC
1501 1550

CAACCCCTGC TGGCCAGGGT CAATGTCTGT CTCTCTGGTC TCTCCAGAAG
1551 1600

TCTTCCATGG CCACCTTCGT CCCCACCCTC CAGAGGAATC TGAAACCGCA
1601 1650

TGTGCTCCCT GGCCCCCACA GCCCCTGCCT CTCCCAGAGC AGCAGTACCT
1651 1700

AAGCCTCAGT GCACTCCAAG AATTGAAACC CTCAGTCTGC TGCCCCTCCC
1701 1750

CACCAGAATG TTTCTCTCCC ATTCTTACCC ACTCAAGGCC CTTTCAGTAG
1751 1800

CCCCTTGGAG TATTCTCTTC CTACATATCA GGGCAACTTC CAAACTCATC
1801 1850

ACCCTTCTGA GGGGTGGGGG AAAGACCCCC ACCACATCGG GGGAGCAGTC

```
1851                                                                1900

CTCCAAGGAC  TGGCCAGTCT  CCAGATGCCC  GTGCACACAG  GAACACTGCC
1901                                                                1950

TTATGCACGG  GAGTCCCAGA  AGAAGGGGTG  ATTTCTTTCC  CCACCTTAGT
1951                                                                2000

TACACCATCA  AGACCCAGCC  AGGGCATCCC  CCCTCCTGGC  CTGAGGGCCA
2001                                                                2050

GCTCCCCATC  CTGAAAAACC  TGTCTGCTCT  CCCCACCCCT  TTGAGGCTAT
2051                                                                2100

AGGGCCCAAG  GGGCAGGTTG  GACTGGATTC  CCCTCCAGCC  CCTCCCGCCC
2101                                                                2150

CCAGGACAAA  ATCAGCCACC  CCAGGGGCAG  GGCCTCACTT  GCCTCAGGAA
2151                                                                2200

CCCCAGCCTG  CCAGCACCTA  TTCCACCTCC  CAGCCCAGCA
2201                                          2239
```

Beginning on page 172 and ending on page 177, the paragraph containing lines 33 and 34 (page 172) has been amended, as follows:

Nucleotide sequence of a mouse uroplakin II 5' flanking region. The translational start site is denoted with an asterisk. SEQ ID NO:[__]7 (number 1 of SEQ ID NO:[6__]7corresponds to position -3592 with respect to the translational start site).

```
 CTCGAGGATCTCGGCCCTCTTTCTGCATCCTTGTCCTAAATCATTTTCAT
1                                                    50

 ATCTTGCTAGACCTCAGTTTGAGAGAAACGAACCTTCTCATTTTCAAGTT
51                                                   100

 GAAAAAAAAAAGAGGTTCAAAGTGGCTCACTCAAAGTTACAAGCCAACAC
101                                                  150

 TCACCACTACGAGTACAATGGCCACCATTAGTGCTGGCATGCCCCAGGAG
151                                                  200

 ACAGGCATGCATATTATTCTAGATGACTGGGAGGCAGAGGGGTGGCCTAG
201                                                  250

 TGAGGTCAGACTGTGGACAGATCAGGCAGATGTGGGTTCTGATCCCAATT
251                                                  300
```

CCTCAGGCCGCAGAACTACTGTGGTTCAAGAAGGGGACAAAAGGACTGCA
301 350

GTCCGGAACAGGAGGTCCATTTGAGAGCTGACTGAGCAGAAGAGGAAAGT
351 400

GAAGAACTTCTGGGGCAAGAGCTTACCCTACTTTACAGCTTTGTTGTCTT
401 450

CTTTACTCCAGGGGCGTCCCTGGTACTCAGTAAATGTCTGTTGGCTTGAG
451 500

GAACATATGTGTAAGGAGGAAGGAGAGGGAACTTGAGGGAGTTAAGACTC
501 550

AAGAATCAATCAAGGAGAGGACAGCAGAGAAGACAGGGTTTGGGAGAGAG
551 600

ACTCCAGACATTGGCCCTGGTTCCCTTCTTGGCCACTGTGAAACCCTCCA
601 650
GAGGAACTGAGTGCTGTGGCTTTAAATGATCTCAGCACTGTCAGTGAAGC
651 700

GCTCTGCTCAAAGAGTTATCCTCTTGCTCCTGTGCCGGGGCCTCCCCCTC
701 750

CTCTCAGCTCCCAAACCCTTCTCAGCCACTGTGATGGCATAATTAGATGC
751 800

GAGAGCTCAGACCGTCAGGTCTGCTCCAGGAACCACCCATTTTCCCCAAC
801 850

CCCAGAGAAAGGTCCTAGTGGAAAAGTGGGGGCCACTGAAGGGCTGATGG
851 900

GGTTCTGTCCTTTCCCCCATGCTGGGTGGACTTAAAGTCTGCGATGTGTG
900 950

TAGGGGGTAGAAGACAACAGAACCTGGGGGCTCCGGCTGGGAGCAGGAGG
951 1000

AACTCTCACCAGACGATCTCCAAATTTACTGTGCAATGGACGATCAGGAA
1001 1050

ACTGGTTCAGATGTAGCTTCTGATACAGTGGGTCTGAGGTAAAACCCGAA
1051 1100

ACTTAATTTCTTTCAAAAATTTAAAGTTGCATTTATTATTTTATATGTGT
1101 1150

GCCCATATGTGTGCCACAGTGTCTATGTGGAGGTCAGAGGGCAAGTTGTG
1151 1200

GGCATTGGCTCTCTCCTTTCATAATGTGGCTTCTGGGGACCAAAATGTCA

1201 1250

GGCATGGTGGCAAGAGCTTTTACCTGTTGAGCCATCTCATGGTTTCGTAA
1251 1300

AACTTCCTATGACGCTTACAGGTAACGCAGAGACACAGACTCACATTTGG
1301 1350

AGTTAGCAGATGCTGTATTGGTGTAAACACTCATACACAGACACACACAC
1351 1400

ATACTCATACACACACACACACTTATCACATGCACACACATACTCGTA
1401 1450

TACACACAGACACACACACATGCACTCTCACATTCACATATTCATACACA
1451 1500

TCCACACACACACTCATCCACACACACAGACACACATACTCATCCACACA
1501 1550

CACACACACACATACTCATACACACACACAGACACACATACTCATACACA
1551 1600

CACACAGACACACACATATAATCATACATACACAGACACACTCATACATG
1601 1650

TGCACACACACACTCATCCACACACACACACTCATACACACACACACTCA
1651 1700

TACACACACACACTCATACACACACACACGAGGTTTTTCTCAGGCTGCCT
1701 1750

TTGGGTGGAGACTGGAACTGATTTCTGTTTTTCAGCTCCTTGGCTTTTTG
1751 1800

TCCCTTTAGATGAGATCTCCTCCTCACTTTACACACAGAAAGATCACACA
1801 1850

CGAGGGAGAACTGGCGGTGCGGAAGAGGGCTACACGGTAGGGTGTCAGGG
1851 1900

TCAGGAGATCTTCCTGGCAAGTCTCAAACCTCCACATAGCACAGTGTTTA
1901 1950

CGTGAGGATTTAGGAGGAATCAGGAAGAGGATTGGTTTACTGCAGAGCAG
1951 2000

ACCATATAGGTCCACTCCTAAGCCCCATTTGAAATTAGAAGTGAGACAGT
2001 2050

GTGGGATAAAAAGAGCAGATCTCTGGTCACATTTTTAAAGGGATATGAGG
2051 3000

GTCCTGTGCCTTTAAGCCTTCCCATCTCCCTCCAATCCCCCCTCACCTTC

2101 2150

CCCACCCTAACCCTCCCCAGGTTTCTGGAGGAGCAGAGTTGCGTCTTCTC
2151 2200

CCTGCCCTGCCGAGCTGCTCACTGGCTGCTCTAGAGGCTGTGCTTTGCGG
2201 2250

TCTCCATGGAAACCATTAGTTGCTAAGCAACTGGAGCATCATCTGTGCTG
2251 2300

AGCTCAGGTCCTATCGAGTTCACCTAGCTGAGACACCCACGCCCCTGCAG
2301 2350

CCACTTTGCAGTGACAAGCCTGAGTCTCAGGTTCTGCATCTATAAAAACG
2351 2400

AGTAGCCTTTCAGGAGGGCATGCAGAGCCCCCTGGCCAGCGTCTAGAGGA
2401 2450

GAGGTGACTGAGTGGGGCCATGTCACTCGTCCATGGCTGGAGAACCTCCA
2451 2500

TCAGTCTCCCAGTTAGCCTGGGGCAGGAGAGAACCAGAGGAGCTGTGGCT
2501 2550

GCTGATTGGATGATTTACGTACCCAATCTGTTGTCCCAGGCATCGAACCC
2551 2600

CAGAGCGACCTGCACACATGCCACCGCTGCCCCGCCCTCCACCTCCTCTG
2601 2650

CTCCTGGTTACAGGATTGTTTTGTCTTGAAGGGTTTTGTTGTTGCTACTT
2651 2700

TTTGCTTTGTTTTTTCTTTTTTAACATAAGGTTTCTCTGTGTAGCCCTAG
2701 2750

CTGTCCTGGAACTCACTCTGTAGACCAGGCTGGCCTCAAACTCAGAAATC
2751 2800

CACCTTCCTCCCAAGTGCTGGGATTAAAGGCATTCGCACCATCGCCCAGC
2801 2850

CCCCGGTCTTGTTTCCTAAGGTTTTCCTGCTTTACTCGCTACCCGTTGCA
2851 2900

CAACCGCTTGCTGTCCAAGTCTGTTTGTATCTACTCCACCGCCCACTAGC
2901 2950

CTTGCTGGACTGGACCTACGTTTACCTGGAAGCCTTCACTAACTTCCCTT
2951 3000

GTCTCCACCTTCTGGAGAAATCTGAAGGCTCACACTGATACCCTCCGCTT

3001 3050

CTCCCAGAGTCGCAGTTTCTTAGGCCTCAGTTAAATACCAGAATTGGATC
3051 3100

TCAGGCTCTGCTATCCCCACCCTACCTAACCAACCCCCTCCTCTCCCATC
3101 3150

CTTACTAGCCAAAGCCCTTTCAACCCTTGGGGCTTTTCCTACACCTACAC
3151 3200

ACCAGGGCAATTTTAGAACTCATGGCTCTCCTAGAAAACGCCTACCTCCT
3201 3250

TGGAGACTGACCCTCTACAGTCCAGGAGGCAGACACTCAGACAGAGGAAC
3251 3300

TCTGTCCTTCAGTCGCGGGAGTTCCAGAAAGAGCCATACTCCCCTGCAGA
3301 3350

GCTAACTAAGCTGCCAGGACCCAGCCAGAGCATCCCCCTTTAGCCGAGGG
3351 3400

CCAGCTCCCCAGAATGAAAAACCTGTCTGGGGCCCCTCCCTGAGGCTACA
3401 3450

GTCGCCAAGGGGCAAGTTGGACTGGATTCCCAGCAGCCCCTCCCACTCCG
3451 3500

AGACAAAATCAGCTACCCTGGGGCAGGCCTCATTGGCCCCAGGAAACCCC
3501 3550
AGCCTGTCAGCACCTGTTCCAGGATCCAGTCCCAGCGCAGTA
3551
3592

On page 177, the paragraph containing line 1 has been amended, as follows:

AFP-TRE. SEQ ID NO:[__]8.

1 GCATTGCTGTGAACTCTGTACTTAGGACTAAACTTTGAGCAATAACACACATAGATTGAG

61 GATTGTTTGCTGTTAGCATACAAACTCTGGTTCAAAGCTCCTCTTTATTGCTTGTCTTGG

121 AAAATTTGCTGTTCTTCATGGTTTCTCTTTTCACTGCTATCTATTTTTCTCAACCACTCA

181 CATGGCTACAATAACTGTCTGCAAGCTTATGATTCCCAAATATCTATCTCTAGCCTCAAT

241 CTTGTTCCAGAAGATAAAAAGTAGTATTCAAATGCACATCAACGTCTCCACTTGGAGGGC

301 TTAAAGACGTTTCAACATACAAACCGGGGAGTTTTGCCTGGAATGTTTCCTAAAATGTGT

361 CCTGTAGCACATAGGGTCCTCTTGTTCCTTAAAATCTAATTACTTTTAGCCCAGTGCTCA

421 TCCCACCTATGGGGAGATGAGAGTGAAAAGGGAGCCTGATTAATAATTACACTAAGTCAA

481 TAGGCATAGAGCCAGGACTGTTTGGGTAAACTGGTCACTTTATCTTAAACTAAATATATC

541 CAAAACTGAACATGTACTTAGTTACTAAGTCTTTGACTTTATCTCATTCATACCACTCAG

601 CTTTATCCAGGCCACTTATGAGCTCTGTGTCCTTGAACATAAAATACAAATAACCGCTAT

661 GCTGTTAATTATTGGCAAATGTCCCATTTTCAACCTAAGGAAATACCATAAAGTAACAGA

721 TATACCAACAAAAGGTTACTAGTTAACAGGCATTGCCTGAAAAGAGTATAAAAGAATTTC

781 AGCATGATTTTCCATATTGTGCTTCCACCACTGCCAATAACA (822)

Beginning on page 177 and ending on page 178, the paragraph containing line 40 (page 177) has been amended, as follows:

Probasin -TRE SEQ ID NO:[ ]9

-426
5'-AAGCTTCCACAAGTGCATTTAGCCTCTCCAGTATTGCTGATGAATCCACAGT

TCAGGTTCAATGGCGTTCAAAACTTGATCAAAAATGACCAGACTTTATATTTA

CACCAACATCTATCTGATTGGAGGAATGGATAATAGTCATCATGTTTAAACAT

CTACCATTCCAGTTAAGAAAATATGATAGCATCTTGTTCTTAGTCTTTTTCTTA
ARE-1

ATAGGGACATAAAGCCCACAAATAAAAATATGCCTGAAGAATGGGACAGGC

ATTGGGCATTGTCCATGCCTAGTAAAGTACTCCAAGAACCTATTTGTATACTA
ARE-2

GATGACACAATGTCAATGTCTGTGTACAACTGCCAACTGGGATGCAAGACAC

TGCCCATG**CCAAT**CATCCTGAAAAGCAGC**TATAAAAA**GCAGGAAGCTACTCT
CAAT box TATAA box

+1 +28
GCACCTTGTCAGTAGGTCCAGATACCTACAG-3'
Transcription site

On page 178, the paragrapgh containing line 6 has been amended, as follows:

Tyrosinase-TRE SEQ ID NO:[ ]10

PinAI end

1 *CCGGT*TGAAAATGATAAGTTGAATTCTGTCTTCGAGAACATAGAAAAGAA

```
51    TTATGAAATGCCAACATGTGGTTACAAGTAATGCAGACCCAAGGCTCCCC
101   AGGGACAAGAAGTCTTGTGTTAACTCTTTGTGGCTCTGAAAGAAAGAGAG
151   AGAGAAAAGATTAAGCCTCCTTGTGGAGATCATGTGATGACTTCCTGATT
201   CCAGCCAGAGCGAGCATTTCCATGGAAACTTCTCTTCCTCTTCACTCGAG
251   ATTACTAACCTTATTGTTAATATTCTAACCATAAGAATTAAACTATTAAT
301   GGTGAATAGAGTTTTTCACTTTAACATAGGCCTATCCCACTGGTGGGATA
351   CGAGCCAATTCGAAAGAAAAAGTCAGTCATGTGCTTTTCAGAGGATGAAA
401   GCTTAAGATAAAGACTAAAAGTGTTTGATGCTGGAGGTGGGAGTGGTATT
451   ATATAGGTCTCAGCCAAGACATGTGATAATCACTGTAGTAGTAGCTGGAA
501   AGAGAAATCTGTGACTCCAATTAGCCAGTTCCTGCAGACCTTGTGA
```

PinAl end

Beginning on page 178 and ending on page 188, the paragraph containing line 20 (page 178) has been amened, as follows:l

Human glandular kallikrein-TRE SEQ ID NO:[__]11

```
gaattcagaa ataggggaag gttgaggaag gacactgaac tcaaagggga tacagtgatt 60
ggtttatttg tcttctcttc acaacattgg tgctggagga attcccaccc tgaggttatg 120
aagatgtctg aacacccaac acatagcact ggagatatga gctcgacaag agtttctcag 180
ccacagagat tcacagccta gggcaggagg acactgtacg ccaggcagaa tgacatggga 240
attgcgctca cgattggctt gaagaagcaa ggactgtggg aggtgggctt tgtagtaaca 300
agagggcagg gtgaactctg attcccatgg gggaatgtga tggtcctgtt acaaattttt 360
caagctggca gggaataaaa cccattacgg tgaggacctg tggagggcgg ctgccccaac 420
tgataaagga aatagccagg tgggggcctt tcccattgta ggggggacat atctggcaat 480
agaagccttt gagacccttt agggtacaag tactgaggca gcaaataaaa tgaaatctta 540
tttttcaact ttatactgca tgggtgtgaa gatatatttg tttctgtaca gggggtgagg 600
gaaaggaggg gaggaggaaa gttcctgcag gtctggtttg gtcttgtgat ccaggggtc 660
ttggaactat ttaaattaaa ttaaattaaa acaagcgact gttttaaatt aaattaaatt 720
aaattaaatt ttactttatt ttatcttaag ttctgggcta catgtgcagg acgtgcagct 780
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ttgttacata ggtaaacgtg tgccatggtg gtttgctgta cctatcaacc catcacctag 840

gtattaagcc cagcatgcat tagctgtttt tcctgacgct ctccctctcc ctgactccca 900

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aaaataggac gaaggtgaaa tattaggtag gaaaagtata atagttgaaa gaagtaaaaa 1260

aaaatatgca tgagtagcag aatgtaaaag aggtgaagaa cgtaatagtg acttttaga 1320

ccagattgaa ggacagagac agaaaaattt taaggaattg ctaaaccatg tgagtgttag 1380

aagtacagtc aataacatta aagcctcagg aggagaaaag aataggaaag gaggaaatat 1440

gtgaataaat agtagagaca tgtttgatgg attttaaaat atttgaaaga cctcacatca 1500

aaggattcat accgtgccat tgaagaggaa gatggaaaag ccaagaagcc agatgaaagt 1560

tagaaatatt attggcaaag cttaaatgtt aaaagtccta gagagaaagg atggcagaaa 1620

tattggcggg aaagaatgca gaacctagaa tataaattca tcccaacagt ttggtagtgt 1680

gcagctgtag ccttttctag ataatacact attgtcatac atcgcttaag cgagtgtaaa 1740

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ggccatgcag tattccatat tgcgtataga tcacattttc tttctttttt ttttttgaga 2460

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aacagtatat aagcattccc ttttctccac agctttgtca tcatggtttt tttttttctt 3060

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caccctgaca gggcaaacag acaacctaca gaatgggagg aaatttttgc aatctattca 3300

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aataatagcc actctgactg gcgtgaaatg gtatctcatt gtggttttca tttgaatttc 3420

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taagaagggg gtgaccaata ggtcagagtc attctggatg caaggggctc cagaggacca 7560

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gcactttcct tgacccccta tctcagatct tgactttgag gttatctcag acttcctcta 7680

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gagaactata aatgtgtatc ctacagggga gaaaaaaaaa aagaactctg aaagagctga 7860

cattttaccg acttgcaaac acataagcta acctgccagt tttgtgctgg tagaactcat 7920

gagactcctg ggtcagaggc aaaagatttt attacccaca gctaaggagg cagcatgaac 7980

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ggtggatgga cacaggggtt tgtggcaaag gtgagcaacc taggcttaga aatcctcaat 8100

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gaggggagga atgtgggttc tgaactctta aatccccaag ggaggagact ggtaaggtcc 8760

cagcttccga ggtactgacg tgggaatggc ctgagaggtc taagaatccc gtatcctcgg 8820

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ccatctactc cttactccac ttgagggtaa tcacctactc ttctagttcc acaagagtgc 9060

gcctgcgcga gtataatctg cacatgtgcc atgtcccgag gcctggggca tcatccactc 9120

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ccttggactc ccaaagcact gggattgctg gcatgagcca ctcaccgtgc ctggcttgca 10860

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ctgtacctgt acgttgtaca ggcatggaga atgaggagtg aggaggccgt accggaaccc 11040

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tctatttcag gaacatggta agttggaggt ccagctctcg ggctcagacg ggtataggga 11280

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tgatctcaag agttggacac cagcctaggc aacatggcca gaccctctct ctacaaaaaa 11520

attaaaaatt agctggatgt ggtggtgcat gcttgtggtc tcagctatcc tggaggctga 11580

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gatcatccca acccctgttg ctgttcatcc tgagcctgcc ttctctggct ttgttcccta 11760

gatcacatct ccatgatcca taggccctgc ccaatctgac ctcacaccgt gggaatgcct 11820

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taggtgccaa cagggcaagg gcggggtcct ggagaatgaa ggctttatag ggctcctcag 12000

ggaggccccc cagccccaaa ctgcaccacc tggccgtgga caccggt 12047

On page 188, the paragraph containing line 37 has been amended, as follows:

HRE-TRE SEQ ID NO:[__]12

ccccgagg cagtgcat gaggctcagg gcgtgcgt gagtcgcagcgagaccccg gggtgcag gccgga

On page 188, the paragraph containing line 42 has been amended, as follows:

PSA-TRE SEQ ID NO:[__]13

aagcttctag ttttcttttc ccggtgacat cgtggaaagc actagcatct ctaagcaatg 60

atctgtgaca atattcacag tgtaatgcca tccagggaac tcaactgagc cttgatgtcc 120

agagattttt gtgttttttt ctgagactga gtctcgctct gtgccaggct ggagtgcagt 180

ggtgcaacct tggctcactg caagctccgc ctcctgggtt cacgccattc tcctgcctca 240

gcctcctgag tagctgggac tacaggcacc cgccaccacg cctggctaat ttttttgtat 300

ttttagtaga gatggggttt cactgtgtta gccaggatgg tctcagtctc ctgacctcgt 360

gatctgccca ccttggcctc ccaaagtgct gggatgacag gcgtgagcca ccgcgcctgg 420

ccgatatcca gagatttttt gggggggctcc atcacacaga catgttgact gtcttcatgg 480

ttgactttta gtatccagcc cctctagaaa tctagctgat atagtgtggc tcaaaacctt 540

cagcacaaat cacaccgtta gactatctgg tgtggcccaa accttcaggt gaacaaaggg 600

actctaatct ggcaggatac tccaaagcat tagagatgac ctcttgcaaa gaaaaagaaa 660

tggaaaagaa aaagaaagaa aggaaaaaaa aaaaaaaaaa gagatgacct ctcaggctct 720

gaggggaaac gcctgaggtc tttgagcaag gtcagtcctc tgttgcacag tctccctcac 780

agggtcattg tgacgatcaa atgtggtcac gtgtatgagg caccagcaca tgcctggctc 840

tggggagtgc cgtgtaagtg tatgcttgca ctgctgaatg gctgggatgt gtcagggatt 900

atcttcagca cttacagatg ctcatctcat cctcacagca tcactatggg atgggtatta 960

ctggcctcat ttgatggaga aagtggctgt ggctcagaaa ggggggacca ctagaccagg 1020

gacactctgg atgctgggga ctccagagac catgaccact caccaactgc agagaaatta 1080

attgtggcct gatgtccctg tcctggagag ggtggaggtg gaccttcact aacctcctac 1140

cttgaccctc tcttttaggg ctctttctga cctccaccat ggtactagga ccccattgta 1200

ttctgtaccc tcttgactct atgaccccca ccgcccactg catccagctg ggtcccctcc 1260

tatctctatt cccagctggc cagtgcagtc tcagtgccca cctgtttgtc agtaactctg 1320

aaggggctga cattttactg acttgcaaac aaataagcta actttccaga gttttgtgaa 1380

tgctggcaga gtccatgaga ctcctgagtc agaggcaaag gcttttactg ctcacagctt 1440

agcagacagc atgaggttca tgttcacatt agtacacctt gccccccccа aatcttgtag 1500

ggtgaccaga gcagtctagg tggatgctgt gcagaagggg tttgtgccac tggtgagaaa 1560

cctgagatta ggaatcctca atcttatact gggacaactt gcaaacctgc tcagcctttg 1620

tctctgatga agatattatc ttcatgatct tggattgaaa acagacctac tctggaggaa 1680

catattgtat cgattgtcct tgacagtaaa caaatctgtt gtaagagaca ttatctttat 1740

tatctaggac agtaagcaag cctggatctg agagagatat catcttgcaa ggatgcctgc 1800

tttacaaaca tccttgaaac aacaatccag aaaaaaaaag gtgttactgt ctttgctcag 1860

aagacacaca gatacgtgac agaaccatgg agaattgcct cccaacgctg ttcagccaga 1920

gccttccacc ctttctgcag gacagtctca acgttccacc attaaatact tcttctatca 1980

catcccgctt ctttatgcct aaccaaggtt ctaggtcccg atcgactgtg tctggcagca 2040

ctccactgcc aaacccagaa taaggcagcg ctcaggatcc cgaaggggca tggctgggga 2100

tcagaacttc tgggtttgag tgaggagtgg gtccaccctc ttgaatttca aaggaggaag 2160

aggctggatg tgaaggtact gggggaggga aagtgtcagt tccgaactct taggtcaatg 2220

agggaggaga ctggtaaggt cccagctccc gaggtactga tgtgggaatg gcctaagaat 2280

ctcatatcct caggaagaag gtgctggaat cctgaggggt agagttctgg gtatatttgt 2340

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gaaggtcccc tgggcaagca caatctgagc atgaaagatg ccccagaggc cttgggtgtc 2580

atccactcat catccagcat cacactctga gggtgtggcc agcaccatga cgtcatgttg 2640

ctgtgactat ccctgcagcg tgcctctcca gccacctgcc aaccgtagag ctgcccatcc 2700

tcctctggtg ggagtggcct gcatggtgcc aggctgaggc ctagtgtcag acagggagcc 2760

tggaatcata gggatccagg actcaaaagt gctagagaat ggccatatgt caccatccat 2820

gaaatctcaa gggcttctgg gtggagggca cagggacctg aacttatggt ttcccaagtc 2880

tattgctctc ccaagtgagt ctcccagata cgaggcactg tgccagcatc agccttatct 2940

ccaccacatc ttgtaaaagg actacccagg gccctgatga acaccatggt gtgtacagga 3000

gtaggggtg gaggcacgga ctcctgtgag gtcacagcca agggagcatc atcatgggtg 3060

gggaggaggc aatggacagg cttgagaacg gggatgtggt tgtatttggt tttctttggt 3120

tagataaagt gctgggtata ggattgagag tggagtatga agaccagtta ggatggagga 3180

tcagattgga gttgggttag ataaagtgct gggtatagga ttgagagtgg agtatgaaga 3240

ccagttagga tggaggatca gattggagtt gggttagaga tggggtaaaa ttgtgctccg 3300

gatgagtttg ggattgacac tgtggaggtg gtttgggatg gcatggcttt gggatggaaa 3360

tagatttgtt ttgatgttgg ctcagacatc cttggggatt gaactgggga tgaagctggg 3420

tttgattttg gaggtagaag acgtggaagt agctgtcaga tttgacagtg gccatgagtt 3480

ttgtttgatg gggaatcaaa caatggggga agacataagg gttggcttgt taggttaagt 3540

tgcgttgggt tgatggggtc ggggctgtgt ataatgcagt tggattggtt tgtattaaat 3600

tgggttgggt caggttttgg ttgaggatga gttgaggata tgcttgggga caccggatcc 3660

atgaggttct cactggagtg gagacaaact tcctttccag gatgaatcca gggaagcctt 3720

aattcacgtg taggggaggt caggccactg gctaagtata tccttccact ccagctctaa 3780

gatggtctta aattgtgatt atctatatcc acttctgtct ccctcactgt gcttggagtt 3840

tacctgatca ctcaactaga aacaggggaa gattttatca aattcttttt ttttttttt 3900

ttttttgag acagagtctc actctgttgc ccaggctgga gtgcagtggc gcagtctcgg 3960

ctcactgcaa cctctgcctc ccaggttcaa gtgattctcc tgcctcagcc tcctgagttg 4020

ctgggattac aggcatgcag caccatgccc agctaatttt tgtattttta gtagagatgg 4080

ggtttcacca atgtttgcca ggctggcctc gaactcctga cctggtgatc cacctgcctc 4140

agcctcccaa agtgctggga ttacaggcgt cagccaccgc gcccagccac ttttgtcaaa 4200

ttcttgagac acagctcggg ctggatcaag tgagctactc tggttttatt gaacagctga 4260

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atctttttt ttttttttt aaatcgaggt ttcagtctca ttctatttcc caggctggag 4440

ttcaatagcg tgatcacagc tcactgtagc cttgaactcc tggccttaag agattctcct 4500

gcttcggtct cccaatagct aagactacag tagtccacca ccatatccag ataattttta 4560

aatttttgg ggggccgggc acagtggctc acgcctgtaa tcccaacacc atgggaggct 4620

gagatgggtg gatcacgagg tcaggagttt gagaccagcc tgaccaacat ggtgaaactc 4680

tgtctctact aaaaaaaaaa aaaatagaaa aattagccgg gcgtggtggc acacggcacc 4740

tgtaatccca gctactgagg aggctgaggc aggagaatca cttgaaccca gaaggcagag 4800

gttgcaatga gccgagattg cgccactgca ctccagcctg ggtgacagag tgagactctg 4860

tctcaaaaaa aaaaaatttt ttttttttt ttgtagagat ggatcttgct ttgtttctct 4920

ggttggcctt gaactcctgg cttcaagtga tcctcctacc ttggcctcgg aaagtgttgg 4980

gattacaggc gtgagccacc atgactgacc tgtcgttaat cttgaggtac ataaacctgg 5040

ctcctaaagg ctaaaggcta aatatttgtt ggagaagggg cattggattt tgcatgagga 5100

tgattctgac ctgggagggc aggtcagcag gcatctctgt tgcacagata gagtgtacag 5160

gtctggagaa caaggagtgg ggggttattg gaattccaca ttgtttgctg cacgttggat 5220

tttgaaatgc tagggaactt tgggagactc atatttctgg gctagaggat ctgtggacca 5280

caagatcttt ttatgatgac agtagcaatg tatctgtgga gctggattct gggttgggag 5340

tgcaaggaaa agaatgtact aaatgccaag acatctattt caggagcatg aggaataaaa 5400

gttctagttt ctggtctcag agtggtgcat ggatcaggga gtctcacaat ctcctgagtg 5460

ctggtgtctt agggcacact gggtcttgga gtgcaaagga tctaggcacg tgaggctttg 5520

tatgaagaat cggggatcgt acccaccccc tgtttctgtt tcatcctggg catgtctcct 5580

ctgcctttgt cccctagatg aagtctccat gagctacaag ggcctggtgc atccagggtg 5640

atctagtaat tgcagaacag caagtgctag ctctccctcc ccttccacag ctctgggtgt 5700

gggagggggt tgtccagcct ccagcagcat ggggagggcc ttggtcagcc tctgggtgcc 5760

agcagggcag gggcggagtc ctggggaatg aaggttttat agggctcctg ggggaggctc 5820

cccagcccca agctt 5835

On page 194, the paragraph has been amended, as follows:

CEA TRE SEQ ID NO: 14

```
aagcttttta gtgctttaga cagtgagctg gtctgtctaa cccaagtgac ctgggctcca      60
tactcagccc cagaagtgaa gggtgaagct gggtggagcc aaaccaggca agcctaccct     120
cagggctccc agtggcctga gaaccattgg acccaggacc cattacttct agggtaagga     180
aggtacaaac accagatcca accatggtct ggggggacag ctgtcaaatg cctaaaaata     240
tacctgggag aggagcaggc aaactatcac tgccccaggt tctctgaaca gaaacagagg     300
ggcaacccaa agtccaaatc caggtgagca ggtgcaccaa atgcccagag atatgacgag     360
gcaagaagtg aaggaaccac ccctgcatca aatgttttgc atgggaagga gaagggggtt     420
gctcatgttc ccaatccagg agaatgcatt tgggatctgc cttcttctca ctccttggtt     480
agcaagacta agcaaccagg actctggatt tggggaaaga cgtttatttg tggaggcaag     540
tgatgacaat cccacgaggg cctaggtgaa gagggcagga aggctcgaga cactggggac     600
tgagtgaaaa ccacacccat gatctgcacc acccatggat gctccttcat tgctcacctt     660
tctgttgata tcagatggcc ccatttttctg taccttcaca gaaggacaca ggctagggtc     720
tgtgcatggc cttcatcccc ggggccatgt gaggacagca ggtgggaaag atcatgggtc     780
ctcctgggtc ctgcagggcc agaacattca tcacccatac tgacctccta gatgggaatg     840
gcttccctgg ggctgggcca acggggcctg ggcaggggag aaaggacgtc aggggacagg     900
gaggaagggt catcgagacc cagcctggaa ggttcttgtc tctgaccatc caggatttac     960
ttccctgcat ctacctttgg tcattttccc tcagcaatga ccagctctgc ttcctgatct    1020
cagcctccca ccctggacac agcacccccag tccctggccc ggctgcatcc acccaatacc    1080
ctgataaccc aggacccatt acttctaggg taaggagggt ccaggagaca gaagctgagg    1140
aaaggtctga agaagtcaca tctgtcctgg ccagaggggga aaaaccatca gatgctgaac    1200
caggagaatg ttgacccagg aaagggaccg aggacccaag aaaggagtca gaccaccagg    1260
gtttgcctga gaggaaggat caaggccccg agggaaagca gggctggctg catgtgcagg    1320
acactggtgg ggcatatgtg tcttagattc tccctgaatt cagtgtccct gccatggcca    1380
gactctctac tcaggcctgg acatgctgaa ataggacaat ggccttgtcc tctctccccca    1440
ccatttggca agagacataa aggacattcc aggacatgcc ttcctgggag gtccaggttc    1500
tctgtctcac acctcaggga ctgtagttac tgcatcagcc atggtaggtg ctgatctcac    1560
ccagcctgtc caggcccttc cactctccac tttgtgacca tgtccaggac caccctcag     1620
atcctgagcc tgcaaatacc cccttgctgg gtgggtggat tcagtaaaca gtgagctcct    1680
```

On page 203, the parargraph containing line 5 has been amended, as follows:

Mucin-TRE SEQ ID NO:[__]15

```
cgagcggccc ctcagcttcg gcgcccagcc ccgcaaggct cccggtgacc actagagggc 60
gggaggagct cctggccagt ggtggagagt ggcaaggaag gaccctaggg ttcatcggag 120
cccaggttta ctcccttaag tggaaatttc ttcccccact cctccttggc tttctccaag 180
gagggaaccc aggctgctgg aaagtccggc tggggcgggg actgtgggtt caggggagaa 240
cggggtgtgg aacgggacag ggagcggtta gaagggtggg gctattccgg gaagtggtgg 300
ggggagggag cccaaaacta gcacctagtc cactcattat ccagccctct tatttctcgg 360
ccgctctgct tcagtggacc cggggagggc ggggaagtgg agtgggagac ctaggggtgg 420
gcttcccgac cttgctgtac aggacctcga cctagctggc tttgttcccc atccccacgt 480
tagttgttgc cctgaggcta aaactagagc ccaggggccc caagttccag actgcccctc 540
ccccctcccc cggagccagg gagtggttgg tgaaaggggg aggccagctg gagaacaaac 600
gggtagtcag ggggttgagc gattagagcc cttgtaccct acccaggaat ggttggggag 660
gaggaggaag aggtaggagg taggggaggg ggcggggttt tgtcacctgt cacctgctcg 720
ctgtgcctag ggcgggcggg cggggagtgg ggggaccggt ataaagcggt aggcgcctgt 780
gcccgctcca cctctcaagc agccagcgcc tgcctgaatc tgttctgccc cctccccacc 840
catttcacca ccaccatg                                              858
```

Beginning on page 203 and ending on page 208, the paragraph containing line 55 (page 203) has been amended, as follows:

αFP-TRE SEQ ID NO:[__]16

```
gaattcttag aaatatgggg gtaggggtgg tggtggtaat tctgttttca ccccataggt 60
gagataagca ttgggttaaa tgtgctttca cacacacatc acatttcata agaattaagg 120
aacagactat gggctggagg actttgagga tgtctgtctc ataacacttg ggttgtatct 180
gttctatggg gcttgtttta agcttggcaa cttgcaacag ggttcactga ctttctcccc 240
aagcccaagg tactgtcctc ttttcatatc tgttttgggg cctctggggc ttgaatatct 300
gagaaaatat aaacatttca ataatgttct gtggtgagat gagtatgaga gatgtgtcat 360
```

tcatttgtat caatgaatga atgaggacaa ttagtgtata aatccttagt acaacaatct 420

gagggtaggg gtggtactat tcaatttcta tttataaaga tacttatttc tatttattta 480

tgcttgtgac aaatgttttg ttcgggacca caggaatcac aaagatgagt ctttgaattt 540

aagaagttaa tggtccagga ataattacat agcttacaaa tgactatgat ataccatcaa 600

acaagaggtt ccatgagaaa ataatctgaa aggtttaata agttgtcaaa ggtgagaggg 660

ctcttctcta gctagagact aatcagaaat acattcaggg ataattattt gaatagacct 720

taagggttgg gtacattttg ttcaagcatt gatggagaag gagagtgaat atttgaaaac 780

attttcaact aaccaaccac ccaatccaac aaacaaaaaa tgaaaagaat ctcagaaaca 840

gtgagataag agaaggaatt ttctcacaac ccacacgtat agctcaactg ctctgaagaa 900

gtatatatct aatatttaac actaacatca tgctaataat gataataatt actgtcattt 960

tttaatgtct ataagtacca ggcatttaga agatattatt ccatttatat atcaaaataa 1020

acttgagggg atagatcatt ttcatgatat atgagaaaaa ttaaaaacag attgaattat 1080

ttgcctgtca tacagctaat aattgaccat aagacaatta gatttaaatt agttttgaat 1140

ctttctaata ccaaagttca gtttactgtt ccatgttgct tctgagtggc ttcacagact 1200

tatgaaaaag taaacggaat cagaattaca tcaatgcaaa agcattgctg tgaactctgt 1260

acttaggact aaactttgag caataacaca catagattga ggattgtttg ctgttagcat 1320

acaaactctg gttcaaagct cctctttatt gcttgtcttg gaaaatttgc tgttcttcat 1380

ggtttctctt ttcactgcta tctatttttc tcaaccactc acatggctac aataactgtc 1440

tgcaagctta tgattcccaa atatctatct ctagcctcaa tcttgttcca gaagataaaa 1500

agtagtattc aaatgcacat caacgtctcc acttggaggg cttaaagacg tttcaacata 1560

caaaccgggg agttttgcct ggaatgtttc ctaaaatgtg tcctgtagca catagggtcc 1620

tcttgttcct taaaatctaa ttacttttag cccagtgctc atcccaccta tggggagatg 1680

agagtgaaaa gggagcctga ttaataatta cactaagtca ataggcatag agccaggact 1740

gtttgggtaa actggtcact ttatcttaaa ctaaatatat ccaaaactga acatgtactt 1800

agttactaag tctttgactt tatctcattc ataccactca gctttatcca ggccacttat 1860

ttgacagtat tattgcgaaa acttcctaac tggtctcctt atcatagtct tatccccttt 1920

tgaaacaaaa gagacagttt caaaatacaa atatgatttt tattagctcc cttttgttgt 1980

ctataatagt cccagaagga gttataaact ccatttaaaa agtctttgag atgtggccct 2040

tgccaacttt gccaggaatt cccaatatct agtattttct actattaaac tttgtgcctc 2100

ttcaaaactg cattttctct cattccctaa gtgtgcattg ttttccctta ccggttggtt 2160

tttccaccac cttttacatt ttcctggaac actataccct ccctcttcat ttggcccacc 2220

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cctccaaaga tgtcatgagt tcctcttttc attctactaa tcacagcatc catcacacca 2340

tgttgtgatt actgatacta ttgtctgttt ctctgattag gcagtaagct caacaagagc 2400

tacatggtgc ctgtctcttg ttgctgatta ttcccatcca aaaacagtgc ctggaatgca 2460

gacttaacat tttattgaat gaataaataa aaccccatct atcgagtgct actttgtgca 2520

agacccggtt ctgaggcatt tatatttatt gatttattta attctcattt aaccatgaag 2580

gaggtactat cactatcctt attttatagt tgataaagat aaagcccaga gaaatgaatt 2640

aactcaccca aagtcatgta gctaagtgac agggcaaaaa ttcaaaccag ttccccaact 2700

ttacgtgatt aatactgtgc tatactgcct ctctgatcat atggcatgga atgcagacat 2760

ctgctccgta aggcagaata tggaaggaga ttggaggatg acacaaaacc agcataatat 2820

cagaggaaaa gtccaaacag gacctgaact gatagaaaag ttgttactcc tggtgtagtc 2880

gcatcgacat cttgatgaac tggtggctga cacaacatac attggcttga tgtgtacata 2940

ttatttgtag ttgtgtgtgt atttttatat atatatttgt aatattgaaa tagtcataat 3000

ttactaaagg cctaccattt gccaggcatt tttacatttg tcccctctaa tcttttgatg 3060

agatgatcag attggattac ttggccttga agatgatata tctacatcta tatctatatc 3120

tatatctata tctatatcta tatctatatc tatatctata tatgtatatc agaaaagctg 3180

aaatatgttt tgtaaagtta taaagatttc agactttata gaatctggga tttgccaaat 3240

gtaacccctt tctctacatt aaacccatgt tggaacaaat acatttatta ttcattcatc 3300

aaatgttgct gagtcctggc tatgaaccag acactgtgaa agcctttggg atattttgcc 3360

catgcttggg caagcttata tagtttgctt cataaaactc tatttcagtt cttcataact 3420

aatacttcat gactattgct tttcaggtat tccttcataa caaatacttt ggctttcata 3480

tatttgagta aagtcccccc tgaggaagag tagaagaact gcactttgta aatactatcc 3540

tggaatccaa acggatagac aaggatggtg ctacctcttt ctggagagta cgtgagcaag 3600

gcctgttttg ttaacatgtt ccttaggaga caaaacttag gagagacacg catagcagaa 3660

aatggacaaa aactaacaaa tgaatgggaa ttgtacttga ttagcattga agaccttgtt 3720

tatactatga taaatgtttg tatttgctgg aagtgctact gacggtaaac ccttttttgtt 3780

taaatgtgtg ccctagtagc ttgcagtatg atctattttt taagtactgt acttagctta 3840

tttaaaaatt ttatgtttaa aattgcatag tgctctttca ttgaagaagt tttgagagag 3900

agatagaatt aaattcactt atcttaccat ctagagaaac ccaatgttaa aactttgttg 3960

tccattattt ctgtctttta ttcaacattt tttttagagg gtgggaggaa tacagaggag 4020

gtacaatgat acacaaatga gagcactctc catgtattgt tttgtcctgt ttttcagtta 4080

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tttgatttta taatcttacc tttagaataa tgaaacacct gtgaagcttt agaaaatact 4260

ggtgcctggg tctcaactcc acagattctg atttaactgg tctgggttac agactaggca 4320

ttgggaattc aaaaagttcc cccagtgatt ctaatgtgta gccaagatcg ggaacccttg 4380

tagacaggga tgataggagg tgagccactc ttagcatcca tcatttagta ttaacatcat 4440

catcttgagt tgctaagtga atgatgcacc tgacccactt tataaagaca catgtgcaaa 4500

taaaattatt ataggacttg gtttattagg gcttgtgctc taagttttct atgttaagcc 4560

atacatcgca tactaaatac tttaaaatgt accttattga catacatatt aagtgaaaag 4620

tgtttctgag ctaaacaatg acagcataat tatcaagcaa tgataatttg aaatgaattt 4680

attattctgc aacttaggga caagtcatct ctctgaattt tttgtacttt gagagtattt 4740

gttatatttg caagatgaag agtctgaatt ggtcagacaa tgtcttgtgt gcctggcata 4800

tgataggcat ttaatagttt taaagaatta atgtatttag atgaattgca taccaaatct 4860

gctgtctttt ctttatggct tcattaactt aatttgagag aaattaatta ttctgcaact 4920

tagggacaag tcatgtcttt gaatattctg tagtttgagg agaatatttg ttatatttgc 4980

aaaataaaat aagtttgcaa gttttttttt tctgccccaa agagctctgt gtccttgaac 5040

ataaaataca aataaccgct atgctgttaa ttattggcaa atgtcccatt ttcaacctaa 5100

ggaaatacca taaagtaaca gatataccaa caaaaggtta ctagttaaca ggcattgcct 5160

gaaaagagta taaaagaatt tcagcatgat tttccatatt gtgcttccac cactgccaat 5220

aaca 5224